SEQUENCE LISTING

	<110>	Li, Haodong Adams, Mark Calenda, Valerie Fataccioli, Virginie											
	<120>	Connective Tissue Growth Factor-2											
	<130>	PF126P2											
	-140-	Imagaianad											
		Unassigned 2001-07-11											
	V1417	2001 07 11											
	<150>	09/348,815											
	<151>	1999-07-08											
	4=0	00/450 303											
	<150> <151>	• •											
	<121>	1995-06-02											
	<150>	> PCT/US94/07736											
	<151>	1994-07-12											
		60/217,402											
	<151>	2000-07-11											
	<150>	60/291,642											
		2001-05-18											
	<160>	8											
	<170>	0> PatentIn version 3.0											
	.0.7.0.	10. 1											
	<210> <211>	1 1146											
		DNA											
		homo sapiens											
	<400>		<i>-</i> 0										
	acgage	coc geategocag ggegetegee ttagtegtea ecetteteea ettgaceagg	60										
	ctggcg	etet ceacetgeee egetgeetge caetgeeeee tggaggegee caagtgegeg	120										
	33 3												
	ccgggag	gtog ggotggtoog ggaoggotgo ggotgotgta aggtotgogo caagoagoto	180										
	22552	70.0t	240										
	aacgag	gact geageaaaac geageeetge gaceaeacea aggggetgga atgeaaette	240										
	ggcgcca	aget ceacegetet gaaggggate tgeagagete agteagaggg cagaceetgt	300										
	gaatata	aact ccagaatcta ccaaaacggg gaaagtttcc agcccaactg taaacatcag	360										
	tgcacai	tgta ttgatggcgc cgtgggctgc attcctctgt gtccccaaga actatctctc	420										
	cccaact	ttgg getgteecaa eeeteggetg gteaaagtta eegggeagtg etgegaggag	480										
	tgggtc	ngtg acgaggatag tatcaaggac cccatggagg accaggacgg cctccttggc	540										
	aaggag	ctgg gattcgatgc ctccgaggtg gagttgacga gaaacaatga attgattgca	600										
Ŧ	attaaaa	aaag gcagctcact gaagcggctc cctgtttttg gaatggagcc tcgcatccta	660										
	2003340												

tacaaccctt tacaaggcca gaaatgtatt gttcaaacaa cttcatggtc ccagtgctca	720									
aagacctgtg gaactggtat ctccacacga gttaccaatg acaaccctga gtgccgcctt	780									
gtgaaagaaa cccggatttg tgaggtgcgg ccttgtggac agccagtgta cagcagcctg	840									
aaaaagggca agaaatgcag caagaccaag aaatcccccg aaccagtcag gtttacttac	900									
gctggatgtt tgagtgtgaa gaaataccgg cccaagtact gcggttcctg cgtggacggc	960									
cgatgctgca cgccccagct gaccaggact gtgaagatgc ggttccgctg cgaagatggg	1020									
gagacatttt ccaagaacgt catgatgatc cagtcctgca aatgcaacta caactgcccg	1080									
catgccaatg aagcagcgtt tcccttctac aggctgttca atgacattca caaatttagg	1140									
gactaa	1146									
<210> 2 <211> 381 <212> PRT <213> homo sapiens										
<400> 2										
Met Ser Ser Arg Ile Ala Arg Ala Leu Ala Leu Val Val Thr Leu Leu 1 5 10 15										
His Leu Thr Arg Leu Ala Leu Ser Thr Cys Pro Ala Ala Cys His Cys 20 25 30										
Pro Leu Glu Ala Pro Lys Cys Ala Pro Gly Val Gly Leu Val Arg Asp 35 40 45										
Gly Cys Gly Cys Cys Lys Val Cys Ala Lys Gln Leu Asn Glu Asp Cys 50 55 60										
Ser Lys Thr Gln Pro Cys Asp His Thr Lys Gly Leu Glu Cys Asn Phe 70 75 80										
Gly Ala Ser Ser Thr Ala Leu Lys Gly Ile Cys Arg Ala Gln Ser Glu 85 90 95										
Gly Arg Pro Cys Glu Tyr Asn Ser Arg Ile Tyr Gln Asn Gly Glu Ser 100 105 110										
Phe Gln Pro Asn Cys Lys His Gln Cys Thr Cys Ile Asp Gly Ala Val 115 120 125										
Gly Cys Ile Pro Leu Cys Pro Gln Glu Leu Ser Leu Pro Asn Leu Gly 130 135 140										
Cys Pro Asn Pro Arg Leu Val Lys Val Thr Gly Gln Cys Cys Glu Glu 145 150 155 160										
Trp Val Cys Asp Glu Asp Ser Ile Lys Asp Pro Met Glu Asp Gln Asp										

Gly Leu Leu Gly Lys Glu Leu Gly Phe Asp Ala Ser Glu Val Glu Leu

Thr	Arg	Asn 195	Asn	Glu	Leu	Ile	Ala 200	Val	Gly	Lys	Gly	Ser 205	Ser	Leu	Lys	
Arg	Leu 210	Pro	Val	Phe	Gly	Met 215	Glu	Pro	Arg	Ile	Leu 220	Tyr	Asn	Pro	Leu	
Gln 225	Gly	Gln	Lys	Cys	Ile 230	Val	Gln	Thr	Thr	Ser 235	Trp	Ser	Gln	Cys	Ser 240	
Lys	Thr	Cys	Gly	Thr 245	Gly	Ile	Ser	Thr	Arg 250	Val	Thr	Asn	Asp	Asn 255	Pro	
Glu	Cys	Arg	Leu 260	Val	Lys	Glu	Thr	Arg 265	Ile	Cys	Glu	Val	Arg 270	Pro	Cys	
Gly	Gln	Pro 275		Tyr	Ser	Ser	Leu 280	Lys	Lys	Gly	Lys	Lys 285	Cys	Ser	Lys	
Thr	Lys 290		Ser	Pro	Glu	Pro 295	Val	Arg	Phe	Thr	Tyr 300	Ala	Gly	Cys	Leu	
Ser		Lys	Lys	Tyr	Arg 310	Pro	Lys	Tyr	Cys	Gly 315	Ser	Cys	Val	Asp	Gly 320	
Arg	Cys	Cys	Thr	Pro 325		Leu	Thr	Arg	Thr 330	Val	Lys	Met	Arg	Phe 335	Arg	
Cys	. Glu	ı Asp	Gly 340		Thr	Phe	Ser	Lys 345	Asn	val	. Met	Met	11e 350	Gln	. Ser	
Cys	s Lys	355		ı Tyr	Asn	Cys	Pro 360	His	Ala	Asr	Glu	Ala 365	Ala	Phe	Pro	
Phe	e Tyr 370		g Lev	ı Phe	e Asr	Asp 375		e His	Lys	s Phe	ar <u>c</u> 380	y Asp)			
	10> 11>	3 28														
	12>	DNA														
<2	13>	hom	o sa	piens	3											
		3														28
cg	cggg	atcc	tgc	gcga	cac a	aatga	agct									20
		4														
	11> 12>															
			o sa	pien	ន											
- 1	00>	4														
		_	caa	tgag	ctc	ccga	atc									27
	10>	5 58														
	112>	DNA	7													
			o sa	pien	ເຮ											
- A	00>	5														
< 4	.00>	ر														

cgctctagat ta	agcgtagt	ctgggacgtc	gtatgggtat	tggaacagcc	tgtagaag	58
<210> 6 <211> 1128 <212> DNA <213> homo s	apiens					
<400> 6						
atgagetece ga	atcgtcag	ggagctcgcc	ttagtcgtca	cccttctcca	cttgaccagg	60
gtggggctct cc	acctgccc	cgctgactgc	cactgccccc	tggaggcgcc	caagtgcgcg	120
ccgggagtcg gg	ctggtccg	ggacggctgc	ggctgttgta	aggtctgcgc	caagcagctc	180
aacgaggact gc	agaaaaac	gcagccctgc	gaccacacca	aggggctgga	atgcaacttc	240
ggcgccagct cc	accgctct	gaaggggatc	tgcagagctc	agtcagaggg	cagaccctgt	300
gaatataact cc	agaatcta	ccaaaacggg	gaaagtttcc	agcccaactg	taaacatcag	360
tgcacatgta tt	ggatggcg	ccggggggct	tgcattcctc	tgtgtcccca	agaactatct	420
ctccccaact tg	ggctgtcc	caaccctcgg	ctggtcaaag	ttaccgggca	gtgctgcgag	480
gagtgggtct gt	gacgagga	tagtatcaag	gaccccatgg	aggaccagga	cggcctcctt	540
ggcaaggggc tg	ggattcga	tgcctccgag	gtggagttga	cgagaaacaa	tgaattgatt	600
gcagttggaa aa	ggcagctc	actgaagcgg	ctccctgttt	ttggaatgga	gcctcgcatc	660
ctatacaacc ct	ttacaagg	ccagaaatgt	attgttcaaa	caacttcatg	gtcccagtgc	720
tcaaagacct gt	ggaactgg	tatctccaca	cgagttacca	atgacaaccc	tgagtgccgc	780
cttgtgaaag aa	acccggat	ttgtgaggtg	cggccttgtg	gacagccagt	gtacagcagc	840
ctgaaaaagg gc	aagaaatg	cagcaagacc	aagaaatccc	ccgaaccagt	caggtttact	900
tacgctggat gt	ttgagtgt	gaagaaatac	cggcccaagt	actgcggttc	ctgcgtggac	960
ggccgatgct gc	acgcccca	gctgaccagg	actgtgaaga	tgcggttccc	ctgcgaagat	1020
ggggagacat tt	tccaagaa	cgtcatgatg	atccagtcct	ccaaatgcaa	ctacaactgc	1080
ccgcatgcca at	gaagcagc	gtttcccttc	tacaggctgt	tccaatga		1128
<210> 7 <211> 375 <212> PRT <213> homo sa	apiens					

<400> 7

Met Ser Ser Arg Ile Val Arg Glu Leu Ala Leu Val Val Thr Leu Leu 1 5 10 15

His Leu Thr Arg Val Gly Leu Ser Thr Cys Pro Ala Asp Cys His Cys 20 25 30

Pro Leu Glu Ala Pro Lys Cys Ala Pro Gly Val Gly Leu Val Arg Asp

40

Pro Phe Tyr Arg Leu Phe Gln 375 370

<210> 8 <211> 30 <212> DNA <213> homo sapiens

<400> 8 cgcgggtacc aggtagcatt tagtccctaa

30